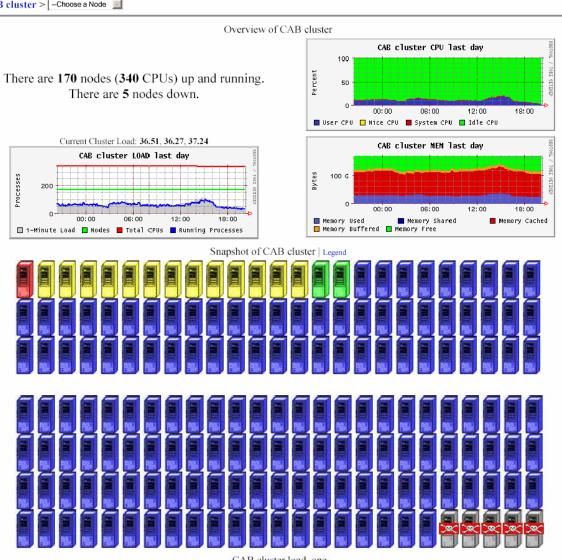
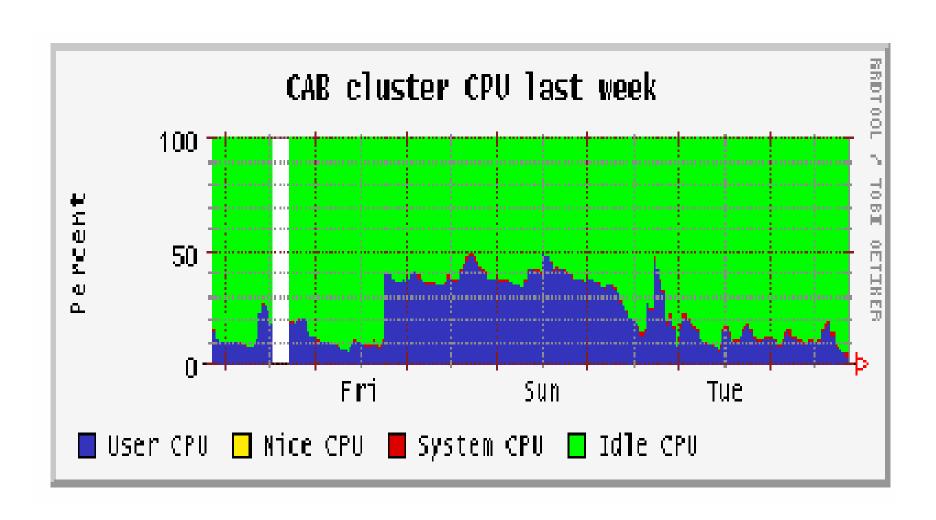
## Cab users report

Dec 19, 2002



 $CPU = 12 \times d0mino$ 5 TB of local scratch for jobs 5 Tb of local cache for sam 100 Mb connections 1 Gb connection to d0mino





#### Access methods

- D0tools (Vivek, Peter...)
  - SAM access fast especially for thumbnails
  - Project areas slow if try to run too many simultaneous jobs.
- mcrunjob (Roger)
- root access not set up yet...

#### documentation

- http://www.nuhep.nwu.edu/~schellma/cab/cab\_doc.html
- http://listserv.fnal.gov/archives/d0cab-users.html
- http://www-clued0.fnal.gov/lancs/cab/index.html
- rund0exe -h

## Sample d0tools command

```
rund0exe -exe=ThumbNailSAM_x -rcp=runThumbNailSAM.rcp \
-rcppkg=thumbnail -localbuild -localrcp -localfwkrcp \
-defname=$definition -maxopt userpostscript=postscript.sh \
-name=$1 \
-cab -scratch=$SCRATCH/junk -jobs=10 -jobname=$1 \
-cabouthost=d0mino.fnal.gov -caboutpath="/prj_root/846/cd_1" \
-cabtar=new.tar
```

- -cab turn it on
- -scratch [–cabtar] path and name of tarball
- [-cabouthost][-caboutpath] other location for output
- [-jobs] = number of parallel jobs
- [-jobname] = name for those jobs

## Parallel jobs

- Thanks to Reiner Hauser, M. Verzocchi and Sam team…
- SAM only
- Sam was designed to run parallel
- You start a project
- It starts a consumer
- Up to 20 consumer processes on 20 machines can attach,
- Your job runs 20 times faster...
- (the 20 is a limit in d0tools, sam on farms runs with 100)

## Known problems

- 1. Can't run large number of jobs off single project disk.
- 2. Sam behavior on timeout is to throw exception which framework does not catch. Do we want to change this?
- 3. If framework gets no files in a parallel job process, it core dumps, things are actually fine as the files went elsewhere but how do we handle this?
- 4. Pbs batch system hangs
  - 1. Bad nodes
  - 2. Heavy traffic (see 1)
- 5. Some code (trigsim?)seems to load in strange libraries from gcc which it shouldn't and can't on the cab...
- 6. Bash not compatible...
- 7. ????

### **Future**

- New d0tools version that has cleaner clued0 submit – M. Verzocchi.
  - setup d0tools –d
    Does this make it possible to submit from bash?
- Make it possible to run from project disk one file at a time by copying to local scratch

## Reprocessing strategies

 On the cab you can run through a lot of data on many nodes fast but that means you will hit a 1/1000 error pretty often.

# For group skims going back into sam as thumbnails

- Get an application family/version
- Say wzskim v00-00-01, bphys-skim v00-00-04
- Check out sam\_manager and change the rcp to contain:

```
string ApplicationName = "wzskip"
string ApplicationVersion = "v00-00-01"
int FileParentageMode=1
string DataTier = "thumbnail-bygroup" // new in test releases
```

Your output will then have metadata with the correct information, identifying the processing as your groups skim with a given version. Store back into sam...

#### Robust resubmission

\_\_set\_\_ yourset minus (file\_analyzed > 0 and data\_tier\_analyzed thumbnail-bygroup and version\_analyzed v00\_00\_00 and appl\_name\_analyzed wzskim)

This is your original input set minus all files that have children from your processing, as opposed to someone elses.

#### Getting only unconsumed files - M. Verzocchi

```
sam create dataset definition --group="dzero" \
    --defname="root1112base" \
    --dim="version p11.12% and data tier root-tuple"
sam create dataset definition --group="dzero"
       --defname="root1112notprocessed" \
    --dim=" set root1112base minus \
   (project name mv-proc1112-% and \
    consumed status consumed and consumer mverzocc )
and then every time you submit a project do
set TIMESTAMP=`date +'%m%d%Y-%H:%M:%S'`
setenv SAM_PROJECT "mv-proc1112-$TIMESTAMP"
sam submit --defname="root1112notprocessed"
```

If a job fails you can recover using the timestamp...